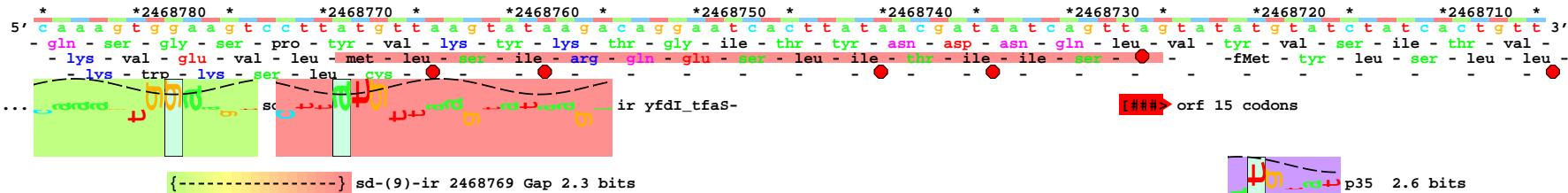
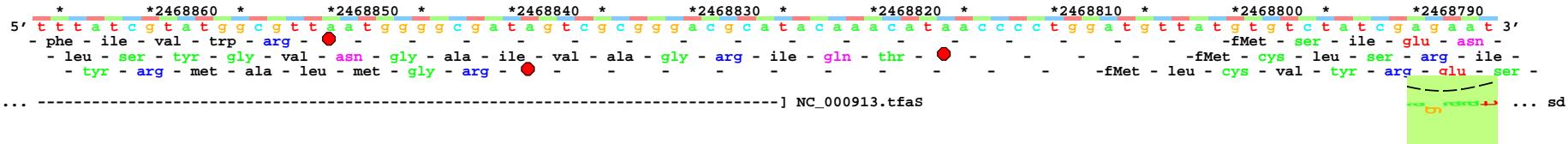
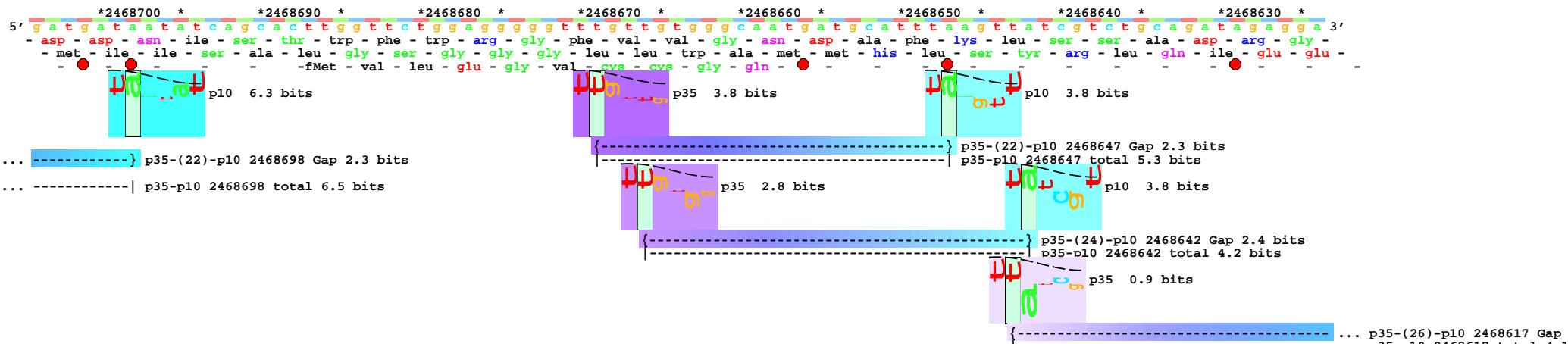


- 1 -

piece 1, NC\_000913, yfdI\_tfaf-, config: linear, direction: -, begin: 2468866, end: 2468465



|-----| sd-ir 2468769 yfdI\_tfaS- total 10.2 bits



Sequence logo showing conservation of amino acids across three positions:

- p10 (6.9 bits)
- p35 (5.6 bits)
- p10 (1.3 bits)

A gap of 2.3 bits is indicated between the second and third positions.

... ----- } p35-(26)-p10 2468617 Gap 3.7 bits  
... ----- | p35-p10 2468617 total 4.1 bits

<sup>5'</sup> \*2468540 \* \*2468530 \* \*2468520 \* \*2468510 \* \*2468500 \* \*2468490 \* \*2468480 \* \*2468470 \* 3'

- gly - glu - ile - arg - leu - val - leu - ● - - fMet - arg - ala - met - ile - gly - ala - gln - phe - lys - leu - ser - ala - leu - his - arg - tyr -

